Docket No.: PF-0162-3 DIV

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By: Many Ramon

Printed: Nancy Ramos

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of: Goli et al.

Title:

A NOVEL GLUTATHIONE S-TRANSFERASE

Serial No.:

To Be Assigned

Filing Date:

Herewith

Examiner:

To Be Assigned

Group Art Unit:

To Be Assigned

Official Draftsman

Commissioner for Patents Washington, D.C. 20231

SUBMISSION OF FORMAL DRAWINGS

Sir:

Transmitted herewith are Figures 1A, 1B, 1C, 2A, 2B, 3, 4, 5, 6, 7, 8, 9 as twelve (12) sheets of formal drawings for this application. Each sheet of drawing indicates the identifying indicia suggested in 37 CFR Section 1.84(c) on the reverse side of the drawings.

Applicants believe that no fee is due with this paper. However, if the Commissioner determines that a fee is necessary, the Commissioner is hereby authorized to charge any additional fees associated with this communication or credit any overpayment to Deposit Account No. **09-0108.** A duplicate copy of this communication is enclosed.

If there are any questions regarding the above, the Examiner is invited to call the undersigned at 650-855-0555.

1

Respectfully submitted,

INCYTE GENOMICS, INC.

Date: 7ebruary 14, 2001

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	TĊC S	108 GAA E	162 CAA Q	216 CTC L	270 ACC T	324 ATG M
54	GAG TCC E S	CTG	CAA Q		AGA R	CTT ATC L I
45	atg	TTT F	TTC F	AGC	GAG	CTT L
	GGC CGG A	99 GAA E	153 CTG L	7 A	.61 .AG	315 2TG
	3 3	GAA E	144 15 AAC CAC CTG CT N H L L	CAG ACC C	AAC CTC N L	306 CTG CTG GAA C L L E I
	AGA R	GAT	CAC H	CAG Q	AAC N	CTG L
36	GGA G	90 TTT F	144 AAC N	198 GTA V	252 14G	306 CTG L
	AAC GGA I	81 9 GA GTC GAG TT	GAT GGT I	189 ATG AAG TTG C M K L	၁၅၅	GAT D
	CCC	GTC V	GAT D	AAG K	TTT F	297 ACA CTG T L
7	CAC TAT	0 0	7	189 ATG M	243 CTC L	297 ACA T
2	CAC H	8CC	TTG L	999	AAT N	999 G
	CTC L	ecc Scc	AAG K	GAC	CAC H	288 GTG GAG V E
တ	CCC AAG O	72 TTA GCT	126 TAC Y	180 ATT I	234 AAG K	288 GTG V
-	P P	TTA L	TTG L	GAA E	GAC	TAC Y
Q	AGG R	GTT V	CAG Q	GTT V	GCA A	ATG M
	ATG GCA GCA AGG	63 TGG W	117 GAA E	171 ATG M	225 AC TAC ATA GCA GAC AA Y I A D K	279 ATT GAC ATG 1 I D M N
	GCA A	AGA R	AAA K	GCC DDD	TAC	ATT I
	ATG M	63 GTG AGA TGG GTT TT V R W V L	ACA T	171 GTG CCC ATG GTT G/ V P M V E	CAC H	CTG L
	5 .					

FIGURE 1A

378	432	486	540	594	548
CAG	GGA	CAA	TTC	CTT	3TC
Q	G	Q	F	L	7
A GCC	CAC	TTA CTC L L	CCT	TTC (648 ACC GTC T V
ATG	GGT	TTA	TTT	AGA	AGA
M		L	F	R	R
369	423	477	531	585	639
AAC	AGG	ATT	GCA	AAG	GTG
N	R	I	A	K	V
GTT	TTA	GTG	TCT	ATT	rat
V	L	V		I	^K
GTG V	ATT I	GAT D	D. J.	ACA	ATT I
360	414	468	522	576	630
GAA	AAG	GCA	ATC	CCT	GAA
E	K	A	I	P	E
AAG	GAA	CTT	AAT	ATC	GAT
K	E		N	I	D
CAA Q	TTT F	AGC S	CCT	AAT N	CCT
351 CAG Q	405 GTG V	459 CTG L	513 ATT I	567 AGT 3	521 2CC
GAT D	CCT	CAG Q	AAA K	AAA CTA Z	CCT P
GAT	TTT	AAT	GAG	AAA	AAG
D	F	N	E	K	K
342	396	450	504	558	612
CCA	TAC	GGT	GAA	GTG	AAG
P	Y	G	E	V	K
AAA	AGA	GTT	CTA	ACA	AAG
K	R	V	L	T	K
TTA L	ATT I	CTT	GCT A	AC	AGC S
333 TTC F	387 ATA I	441 TTT F	495 TTA L	549 GAA	603 r GGC G
CCT	387 AAG GCT ATA ATT AG C A I I R	AGC	ATT I	CAG Q	CCT
333 CAT CCT TTC T H P F L	AAG K	441 CAA AGC TTT C Q S F L	495 ACC ATT TTA G T I L A	CTC	GAA E

FIGURE 1B

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	AAC	Z
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756	ATG
	GTC
	\mathtt{TCT}
747	AGC
	သသ
	GAT
738	ATG
	TTA
	GTC
729	CAT
	AGT
	TAC
720	GTC
	ATT
	GGT
711	GAT
	AGA
	CCT

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792	ָרָרָי הַלָּי
	C.Tr.T.
	TAA
783	TCC TAA
	999
	GTT
774	TAA
	TAT
	ATG
765	\mathbf{TCT}
	CTA
	GTG

FIGURE 1C

g259141 g259141 SAED FLETKEQ SAED LYKLODGNHLLFQQVPMVEIDGMKLVQTRSILHYIADKHN LDKLRNDGYLMFQQVPMVEIDGMKLVQTRAILNYIASKYN LDKLRNDGYLMFQQVPMVEIDGMKLVQTRAILNYIASKYN YEKMORDGHLLFGQVPLVEIDGMMLTQTRAILSYLAAKYN FGKNLKERTLIDMYVEGTLDLLELLIMHPFLKPDDQQKE YGKDIKEKALIDMYIEGIADLGEMILLLPFTQPEEQDAK YGKDIKERALIDMYIEGIADLGEMILLLPVCPPEEKDAK YGKDLKERVRIDMYADGTQDLMMMIAVAPFKTPKEKES V V N M A Q K A I I R Y F P V F E K I L R G H G Q S F L V G N Q L S L A D V I L L A L I Q E K T K N R Y F P A F E K V L K S H G Q D Y L V G N K L S R A D I H L L A L I K E K I K N R Y F P A F E K V L K S H G Q D Y L V G N K L S R A D I H L Y D L I L S R A K T R Y F P V F E K I L K D H G E A F L V G N Q L S W A D I Q L [1] TR VEFD MAARPKLHYPNGRGRMESVRWVLAAAGVE MAEKPKLHYSNTRGRMESIRWLLAAAGVE MAEKPKLHYFNARGRMESTRWLLAAAGVE MAAKPKLYYFNGRGRMESIRWLLAAAGVE ΓY 44 ਲ ਲ ਲ ਲ 44

FIGURE 2A

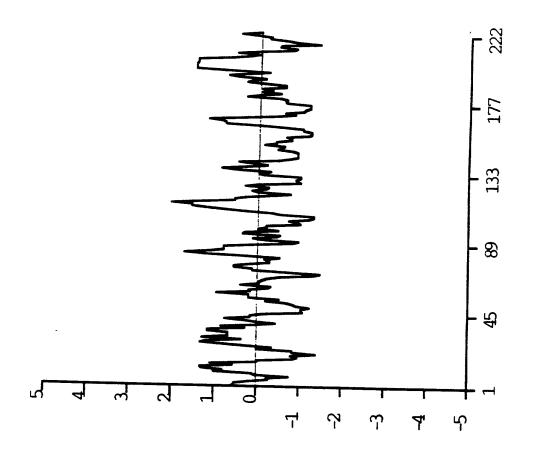
g825605 g259141

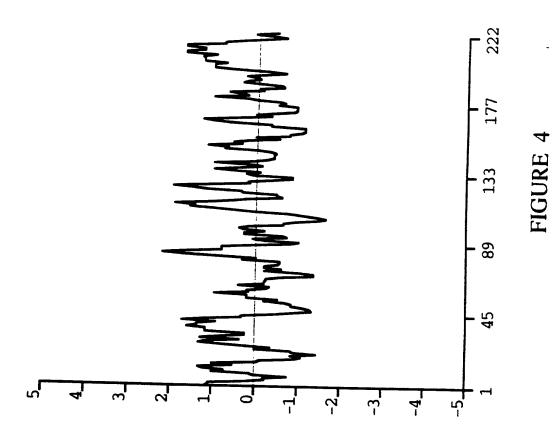
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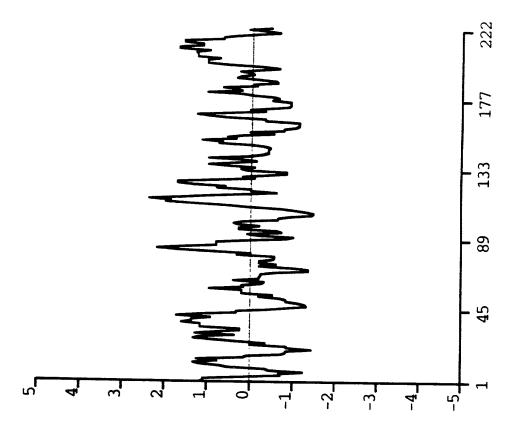
HCST g825605 g259141 q193710
0 0 0 0
PTIKI, PTVKI
KLSNI RISNI RISNI RISNI
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K I P N I L S A F P F L O I L D S S L I S S F P L L K L D S S L I S S F P L L K L S A P V L S D F P L L O
PNILS SSLIS SSLIS SSLIS APVLS
1 1
L Q T I L A L E V E L L Y Y V E V E L L Y Y V E L E A I L M V E
161 L Q 161 V E 161 V E 161 L E
H

201 G SKKKP PPD EIYUR TUYNIFRP 201 G S P R K P P M D E K S L E E S R K I F R F 201 G S P R K P P M D E K S L E E A R K I F R F 201 G S OR K P P D G P Y V E V V R I V L K F

FIGURE 2B







igure 5

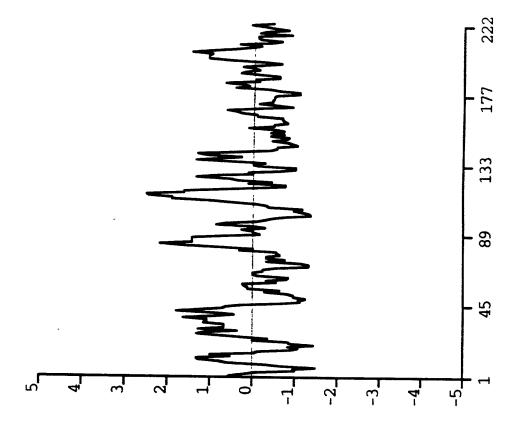


FIGURE 6

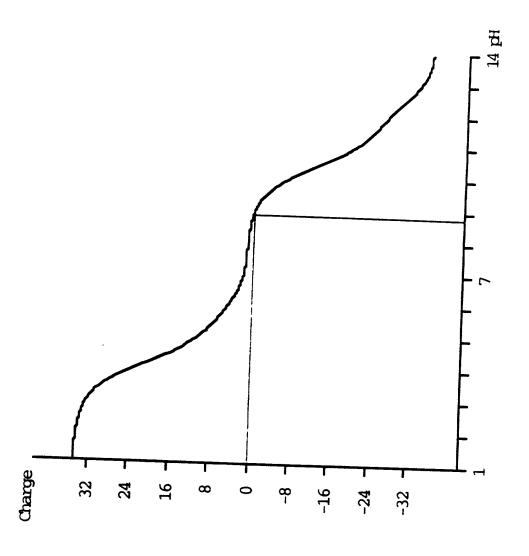


FIGURE 7

